



1962-4066.ST25.txt



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SEQUENCE LISTING

<110> Hoechst Marion Russel  
KATSUURA, MIEKO  
KIMURA, MICHIO

<120> BONE MORPHOGENETIC PROTEIN ANTAGONIST BASED ON THE MATURE  
PROTEIN

<130> 447.001

<140> US 09/806,368

<141> 2001-03-28

<150> PCT/IB99/01621

<151> 1999-10-04

<150> JP 10/288,103

<151> 1998-10-09

<160> 7

<170> PatentIn version 3.1

<210> 1

<211> 119

<212> PRT

<213> Homo sapiens

<220>

<221> CHAIN

<222> (1)..(119)

<223> Mature MP52

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<302> NOVEL PROTEIN AND PROCESS FOR PRODUCING THE SAME

<309>

<310> WO9633215

<311> 1996-04-19

<312> 1996-10-24

<313> (1)..(119)

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Pro Leu Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala  
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Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp  
20 25 30

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu  
35 40 45

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His  
50 55 60

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro  
65 70 75 80

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe  
85 90 95

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val  
100 105 110

Val Glu Ser Cys Gly Cys Arg  
115

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<310> W08800205  
<311> 1987-06-30  
<312> 1988-01-14  
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His Pro Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn Asp Trp Ile  
20 25 30

Val Ala Pro Pro Gly Tyr His Ala Phe Tyr Cys His Gly Glu Cys Pro  
35 40 45

Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala Ile Val Gln  
50 55 60

Thr Leu Val Asn Ser Val Asn Ser Lys Ile Pro Lys Ala Cys Cys Val  
65 70 75 80

Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp Glu Asn Glu  
                   85                  90                  95

Lys Val Val Leu Lys Asn Tyr Gln Asp Met Val Val Glu Gly Cys Gly  
                   100                  105                  110

Cys Arg

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<300>  
 <301> Wozney, JM et al.  
 <302> NOVEL REGULATORS OF BONE FORMATION MOLECULAR CLONES AND  
 ACTIVITIES  
 <303> SCIENCE  
 <304> 242  
 <305> 4885  
 <306> 1528-1534  
 <307> 1988-12-16  
 <308> GENBANK/M22490  
 <309> 1994-10-31

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Ser Pro Lys His His Ser Gln Arg Ala Arg Lys Lys Asn Lys Asn Cys  
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Arg Arg His Ser Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn Asp  
                   20                  25                  30

Trp Ile Val Ala Pro Pro Gly Tyr Gln Ala Phe Tyr Cys His Gly Asp  
                   35                  40                  45

Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala Ile  
                   50                  55                  60

Val Gln Thr Leu Val Asn Ser Val Asn Ser Ser Ile Pro Lys Ala Cys  
 65                  70                  75                  80

Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp Glu  
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Tyr Asp Lys Val Val Leu Lys Asn Tyr Gln Glu Met Val Val Glu Gly  
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Cys Gly Cys Arg  
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<300>  
 <301> OZKAYNAK, E. et al.  
 <302> OP-1 cDNA encodes an osteogenic protein in the TGF-beta.  
 <303> EMBO J.  
 <304> 9  
 <305> 7  
 <306> 2085-2093  
 <307> 1990-07-01  
 <308> EMBL/ X51801  
 <309> 1994-10-31

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Ser Thr Gly Ser Lys Gln Arg Ser Gln Asn Arg Ser Lys Thr Pro Lys  
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Asn Gln Glu Ala Leu Arg Met Ala Asn Val Ala Glu Asn Ser Ser Ser  
 20 25 30

Asp Gln Arg Gln Ala Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg  
 35 40 45

Asp Leu Gly Trp Gln Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala  
 50 55 60

Tyr Tyr Cys Glu Gly Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn  
 65 70 75 80

Ala Thr Asn His Ala Ile Val Gln Thr Leu Val His Phe Ile Asn Pro  
 85 90 95

Glu Thr Val Pro Lys Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile

100                      105                      110  
 Ser Val Leu Tyr Phe Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr  
           115                      120                      125  
  
 Arg Asn Met Val Val Arg Ala Cys Gly Cys His  
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 are modified to Met sulfoxide.  
  
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 Pro Leu Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala  
 1                      5                      10                      15  
  
 Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp  
           20                      25                      30  
  
 Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu  
           35                      40                      45  
  
 Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His  
           50                      55                      60  
  
 Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro  
 65                      70                      75                      80  
  
 Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe  
           85                      90                      95  
  
 Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val  
           100                      105                      110  
  
 Val Glu Ser Cys Gly Cys Arg  
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<212> PRT  
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<220>  
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 and/or 111th Met are modified to s-carboxymethyl Met.

<220>  
 <221> CHAIN  
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 <223> Mature MP52 protein. Note : 30th and/or 71st and/or 74th  
 and/or 111th Met are modified to s-carboxymethyl Met.

<400> 6

Pro Leu Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala  
 1 5 10 15

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp  
 20 25 30

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu  
 35 40 45

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His  
 50 55 60

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro  
 65 70 75 80

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe  
 85 90 95

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val  
 100 105 110

Val Glu Ser Cys Gly Cys Arg  
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<210> 7  
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 <212> PRT  
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<220>  
 <221> CHAIN  
 <222> (1)..(119)  
 <223> Mature MP52 protein. Note : 32nd and 35th Trp are modified  
 to allylsulphenyl Trp.

&lt;400&gt; 7

Pro Leu Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala  
 1 5 10 15

Arg Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp  
 20 25 30

Asp Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu  
 35 40 45

Gly Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His  
 50 55 60

Ala Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro  
 65 70 75 80

Pro Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe  
 85 90 95

Ile Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val  
 100 105 110

Val Glu Ser Cys Gly Cys Arg  
 115